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09/13

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OIEP

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/945,265

DATE: 09/21/2001
 TIME: 20:49:59

Input Set : A:\Seqlist.txt
 Output Set: N:\CRF3\09212001\I945265.raw

ENTERED

3 <110> APPLICANT: Springer, Timothy A.
 4 Shimaoka, Motomu
 5 Lu, Chafen
 7 <120> TITLE OF INVENTION: MODIFIED POLYPEPTIDES STABILIZED IN A
 8 DESIRED CONFORMATION AND METHODS FOR PRODUCING SAME
 10 <130> FILE REFERENCE: CBN-002CP
 C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/945,265
 C--> 12 <141> CURRENT FILING DATE: 2001-08-31
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 13 <151> PRIOR FILING DATE: 2000-09-01
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 35 atttcagaaa attctggact tcatgaagga tgtgatgaag aaactcagca acacttcgta 660
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56 tgaaggggat ggcttggcag atgtggctgt gggggctgag agccagatga tcgtgctgag 1920
57 ctcccggccc gtggtggata tggtcaccct gatgtccttc tctccagctg agatcccagt 1980
58 gcatgaagtg gagtgcctct attcaaccag taacaagatg aaagaaggag ttaatatcac 2040
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99 acgtacttag cagctatctc tcagtgaact gtgagggtaa aggtataact tgtcttgttc 4500
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103 tgaggtcagg agttagaggc cagcctggcg aaaccccatc tctactaaaa atacaaaatc 4740
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122 Ala Arg Ser Phe Ser Pro Pro Arg Ala Gly Arg His Phe Gly Tyr Arg
123 35 40 45
124 Val Leu Gln Val Gly Asn Gly Val Ile Val Gly Ala Pro Gly Glu Gly
125 50 55 60
126 Asn Ser Thr Gly Ser Leu Tyr Gln Cys Gln Ser Gly Thr Gly His Cys
127 65 70 75 80
128 Leu Pro Val Thr Leu Arg Gly Ser Asn Tyr Thr Ser Lys Tyr Leu Gly
129 85 90 95
130 Met Thr Leu Ala Thr Asp Pro Thr Asp Gly Ser Ile Leu Ala Cys Asp
131 100 105 110
132 Pro Gly Leu Ser Arg Thr Cys Asp Gln Asn Thr Tyr Leu Ser Gly Leu
133 115 120 125
134 Cys Tyr Leu Phe Arg Gln Asn Leu Gln Gly Pro Met Leu Gln Gly Arg
135 130 135 140
136 Pro Gly Phe Gln Glu Cys Ile Lys Gly Asn Val Asp Leu Val Phe Leu
137 145 150 155 160
138 Phe Asp Gly Ser Met Ser Leu Gln Pro Asp Glu Phe Gln Lys Ile Leu
139 165 170 175
140 Asp Phe Met Lys Asp Val Met Lys Lys Leu Ser Asn Thr Ser Tyr Gln
141 180 185 190
142 Phe Ala Ala Val Gln Phe Ser Thr Ser Tyr Lys Thr Glu Phe Asp Phe
143 195 200 205
144 Ser Asp Tyr Val Lys Arg Lys Asp Pro Asp Ala Leu Leu Lys His Val
145 210 215 220
146 Lys His Met Leu Leu Leu Thr Asn Thr Phe Gly Ala Ile Asn Tyr Val
147 225 230 235 240
148 Ala Thr Glu Val Phe Arg Glu Glu Leu Gly Ala Arg Pro Asp Ala Thr
149 245 250 255
150 Lys Val Leu Ile Ile Ile Thr Asp Gly Glu Ala Thr Asp Ser Gly Asn
151 260 265 270
152 Ile Asp Ala Ala Lys Asp Ile Ile Arg Tyr Ile Ile Gly Ile Gly Lys
153 275 280 285
154 His Phe Gln Thr Lys Glu Ser Gln Glu Thr Leu His Lys Phe Ala Ser

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156 Lys Pro Ala Ser Glu Phe Val Lys Ile Leu Asp Thr Phe Glu Lys Leu
157 305      310      315      320
158 Lys Asp Leu Phe Thr Glu Leu Gln Lys Lys Ile Tyr Val Ile Glu Gly
159      325      330      335
160 Thr Ser Lys Gln Asp Leu Thr Ser Phe Asn Met Glu Leu Ser Ser Ser
161      340      345      350
162 Gly Ile Ser Ala Asp Leu Ser Arg Gly His Ala Val Val Gly Ala Val
163      355      360      365
164 Gly Ala Lys Asp Trp Ala Gly Gly Phe Leu Asp Leu Lys Ala Asp Leu
165      370      375      380
166 Gln Asp Asp Thr Phe Ile Gly Asn Glu Pro Leu Thr Pro Glu Val Arg
167 385      390      395      400
168 Ala Gly Tyr Leu Gly Tyr Thr Val Thr Trp Leu Pro Ser Arg Gln Lys
169      405      410      415
170 Thr Ser Leu Leu Ala Ser Gly Ala Pro Arg Tyr Gln His Met Gly Arg
171      420      425      430
172 Val Leu Leu Phe Gln Glu Pro Gln Gly Gly Gly His Trp Ser Gln Val
173      435      440      445
174 Gln Thr Ile His Gly Thr Gln Ile Gly Ser Tyr Phe Gly Gly Glu Leu
175      450      455      460
176 Cys Gly Val Asp Val Asp Gln Asp Gly Glu Thr Glu Leu Leu Ile
177-465      470      475      480
178 Gly Ala Pro Leu Phe Tyr Gly Glu Gln Arg Gly Gly Arg Val Phe Ile
179      485      490      495
180 Tyr Gln Arg Arg Gln Leu Gly Phe Glu Glu Val Ser Glu Leu Gln Gly
181      500      505      510
182 Asp Pro Gly Tyr Pro Leu Gly Arg Phe Gly Glu Ala Ile Thr Ala Leu
183      515      520      525
184 Thr Asp Ile Asn Gly Asp Gly Leu Val Asp Val Ala Val Gly Ala Pro
185      530      535      540
186 Leu Glu Glu Gln Gly Ala Val Tyr Ile Phe Asn Gly Arg His Gly Gly
187 545      550      555      560
188 Leu Ser Pro Gln Pro Ser Gln Arg Ile Glu Gly Thr Gln Val Leu Ser
189      565      570      575
190 Gly Ile Gln Trp Phe Gly Arg Ser Ile His Gly Val Lys Asp Leu Glu
191      580      585      590
192 Gly Asp Gly Leu Ala Asp Val Ala Val Gly Ala Glu Ser Gln Met Ile
193      595      600      605
194 Val Leu Ser Ser Arg Pro Val Val Asp Met Val Thr Leu Met Ser Phe
195      610      615      620
196 Ser Pro Ala Glu Ile Pro Val His Glu Val Glu Cys Ser Tyr Ser Thr
197 625      630      635      640
198 Ser Asn Lys Met Lys Glu Gly Val Asn Ile Thr Ile Cys Phe Gln Ile
199      645      650      655
200 Lys Ser Leu Tyr Pro Gln Phe Gln Gly Arg Leu Val Ala Asn Leu Thr
201      660      665      670
202 Tyr Thr Leu Gln Leu Asp Gly His Arg Thr Arg Arg Gly Leu Phe
203      675      680      685

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204 Pro Gly Gly Arg His Glu Leu Arg Arg Asn Ile Ala Val Thr Thr Ser
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207 705      710      715      720
208 Leu Ile Ser Pro Ile Asn Val Ser Leu Asn Phe Ser Leu Trp Glu Glu
209      725      730      735
210 Glu Gly Thr Pro Arg Asp Gln Arg Ala Gln Gly Lys Asp Ile Pro Pro
211      740      745      750
212 Ile Leu Arg Pro Ser Leu His Ser Glu Thr Trp Glu Ile Pro Phe Glu
213      755      760      765
214 Lys Asn Cys Gly Glu Asp Lys Lys Cys Glu Ala Asn Leu Arg Val Ser
215      770      775      780
216 Phe Ser Pro Ala Arg Ser Arg Ala Leu Arg Leu Thr Ala Phe Ala Ser
217 785      790      795      800
218 Leu Ser Val Glu Leu Ser Leu Ser Asn Leu Glu Glu Asp Ala Tyr Trp
219      805      810      815
220 Val Gln Leu Asp Leu His Phe Pro Pro Gly Leu Ser Phe Arg Lys Val
221      820      825      830
222 Glu Met Leu Lys Pro His Ser Gln Ile Pro Val Ser Cys Glu Glu Leu
223      835      840      845
224 Pro Glu Glu Ser Arg Leu Leu Ser Arg Ala Leu Ser Cys Asn Val Ser
225      850      855      860
226 Ser Pro Ile Phe Lys Ala Gly His Ser Val Ala Leu Gln Met Met Phe
227 865      870      875      880
228 Asn Thr Leu Val Asn Ser Ser Trp Gly Asp Ser Val Glu Leu His Ala
229      885      890      895
230 Asn Val Thr Cys Asn Asn Glu Asp Ser Asp Leu Leu Glu Asp Asn Ser
231      900      905      910
232 Ala Thr Thr Ile Ile Pro Ile Leu Tyr Pro Ile Asn Ile Leu Ile Gln
233      915      920      925
234 Asp Gln Glu Asp Ser Thr Leu Tyr Val Ser Phe Thr Pro Lys Gly Pro
235      930      935      940
236 Lys Ile His Gln Val Lys His Met Tyr Gln Val Arg Ile Gln Pro Ser
237 945      950      955      960
238 Ile His Asp His Asn Ile Pro Thr Leu Glu Ala Val Val Gly Val Pro
239      965      970      975
240 Gln Pro Pro Ser Glu Gly Pro Ile Thr His Gln Trp Ser Val Gln Met
241      980      985      990
242 Glu Pro Pro Val Pro Cys His Tyr Glu Asp Leu Glu Arg Leu Pro Asp
243      995      1000      1005
244 Ala Ala Glu Pro Cys Leu Pro Gly Ala Leu Phe Arg Cys Pro Val Val
245      1010      1015      1020
246 Phe Arg Gln Glu Ile Leu Val Gln Val Ile Gly Thr Leu Glu Leu Val
247 1025      1030      1035      1040
248 Gly Glu Ile Glu Ala Ser Ser Met Phe Ser Leu Cys Ser Ser Leu Ser
249      1045      1050      1055
250 Ile Ser Phe Asn Ser Ser Lys His Phe His Leu Tyr Gly Ser Asn Ala
251      1060      1065      1070
252 Ser Leu Ala Gln Val Val Met Lys Val Asp Val Val Tyr Glu Lys Gln

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VERIFICATION SUMMARY

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L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date